

10/533054

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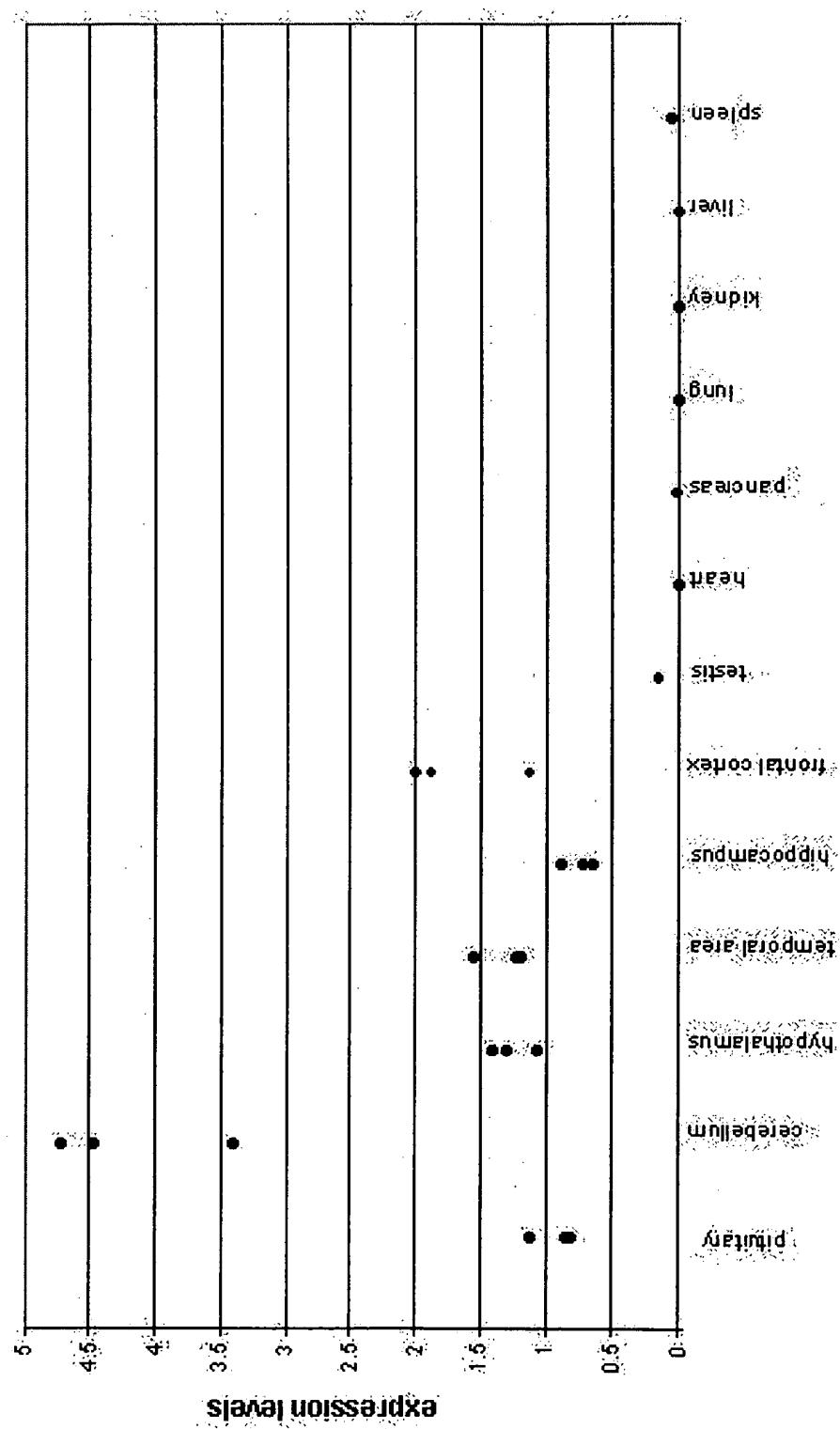
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Table 1

Gene Name	CDS	SEQ ID	NA	AA	Title	Sequence Derived from
down	1700016E05Rik	1	RIKEN cDNA 1700016E05 gene			AB22377
down	23000210BRK	2	RIKEN cDNA 23000210 gene			AB22384
down	2810008D04Rik	3	RIKEN cDNA 2810008D04 gene			AB22387
down	2810511602Rik	4	RIKEN cDNA 2810511602 gene			AB221884
down	4831430D01Rik	5	RIKEN cDNA 4831430D1 gene			AB26942
down	AI414047	6	expressed sequence AI414047			AB2125649
down	Genetra:47654	7	ESTs, moderately similar to V153: MOUSE:VISININ-LIKE PROTEIN 3 [M: musculus]			CB0148
down		8	ESTs, moderately similar to V153: MOUSE: VISININ-LIKE PROTEIN 3 [M: musculus]			AB43393
down	B-F-ABP	9	Brain Fatty acid-binding protein gene			U04827
down	B-F-ABP	10	Brain Fatty acid-binding protein			AA4481304
down	Edg2	214:1308	11	12	endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor 2	U13370
down	Fam2	1:1038	13	14	fibroblast growth factor receptor 2	M22382
down	Gab1	24:2111	15	16	growth factor receptor bound protein 2-associated protein	AB08826
down	Hpcal1	26:1842	17	18	hippocalcin-like 1	AF085182
down	Hsd11b1	128:1006	19	20	hydroxysteroid 11-beta dehydrogenase 1	X83202
down	Nis12	27:1280	21	22	neurotensin receptor 2	U51908
down	Nis11	244:1718	40	41	neurotensin receptor 1	NM_01876
down	Ywhaq	21:758	23	24	tyrosine 3-monooxygenase, tryptophan 5-monooxygenase activation protein, theta polypeptide	U58243
up	1110014Q22Rik	25			RIKEN cDNA 1110014Q22 gene	AB22646
up	1200008D14Rik	26			RIKEN cDNA 1200008D14 gene	AB22638
up	1600023A02Rik	27			RIKEN cDNA 1600023A02 gene	AB219397
up	3110003A17Rik	28			RIKEN cDNA 3110003A17 gene	AB233425
up	3110023110Rik	29			RIKEN cDNA 3110023110 gene	AB196392
up	AI407667	30			expressed sequence AI407667	AB53024
up	AI480570	31			expressed sequence AI480570	AB40646
up	Fkbp5	220:1690	32	33	Fkbp5 binding protein 5 (51 kDa)	U11689
up	Pdyn	1:747	34	35	prodynorphin	AB382264
up	Pank1	287:1408	36	37	preproenkephalin 1	MS9181
up	Sak	69:1384	38	39	serum glucocorticoid regulated kinase 1	AB046181

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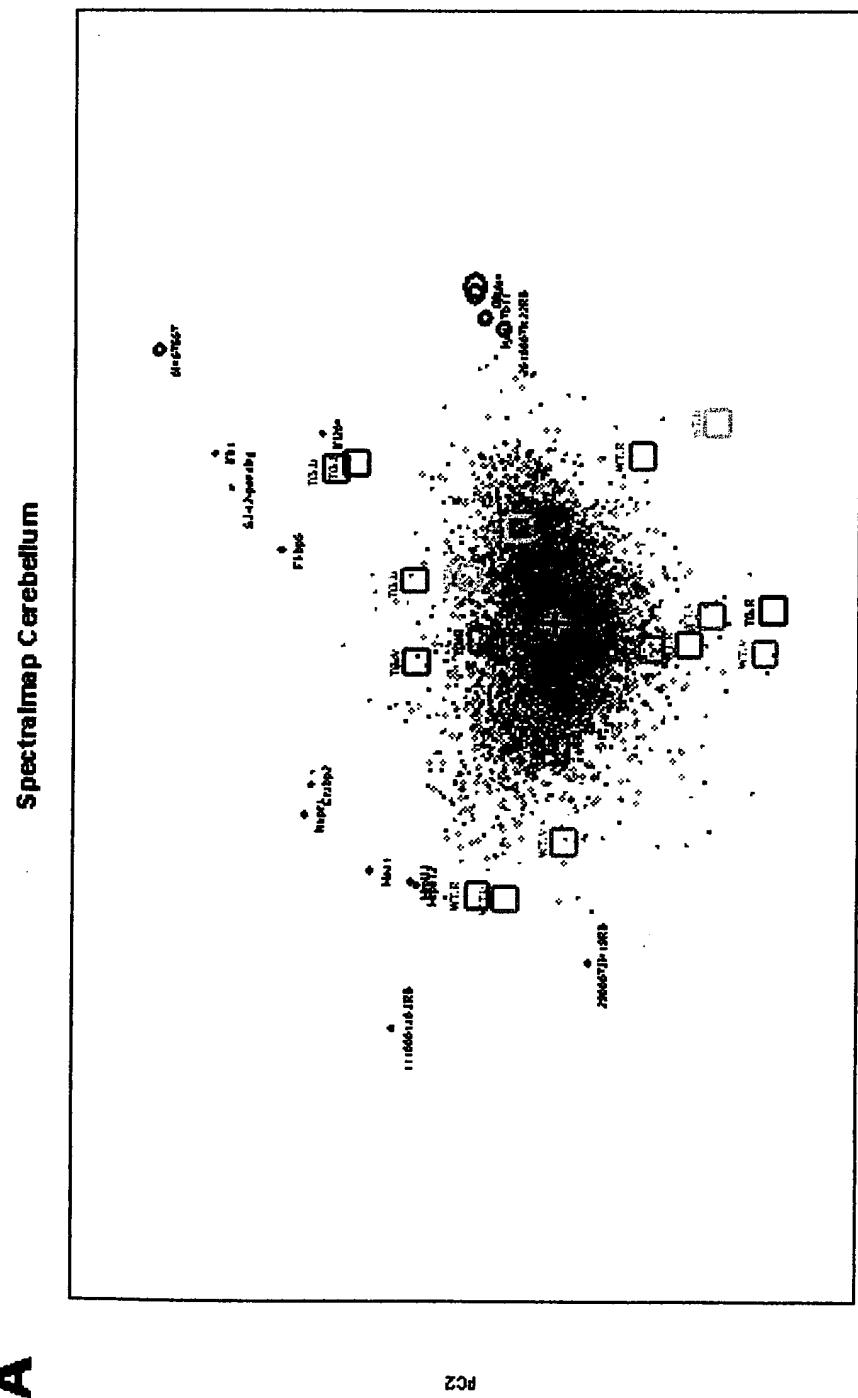
Fig. 1



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Fig. 2

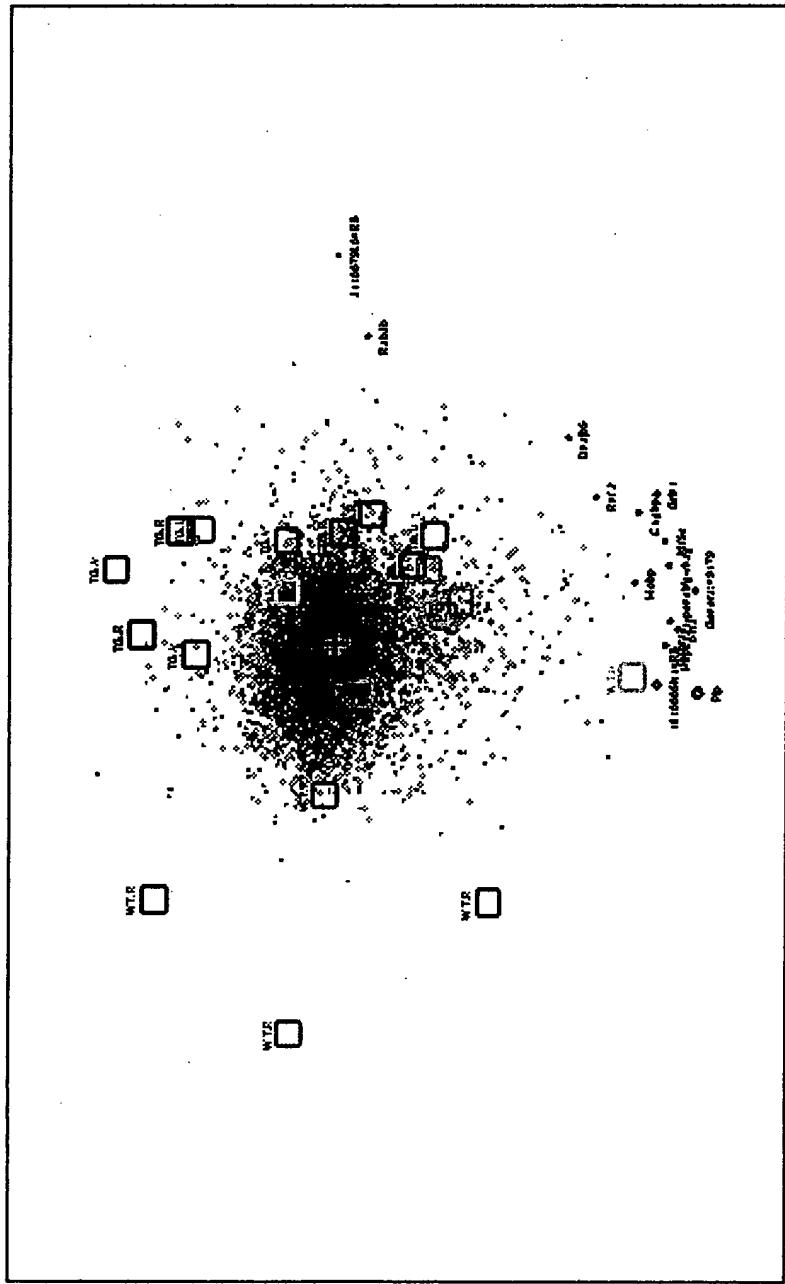


Repres. = 47 % + 12 %, Closure = none, Center = double, Norm. = global, Scale = unc, RUU = mean, C00 = mean

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Fig.2 continued

Spectralmap Frontal Cortex

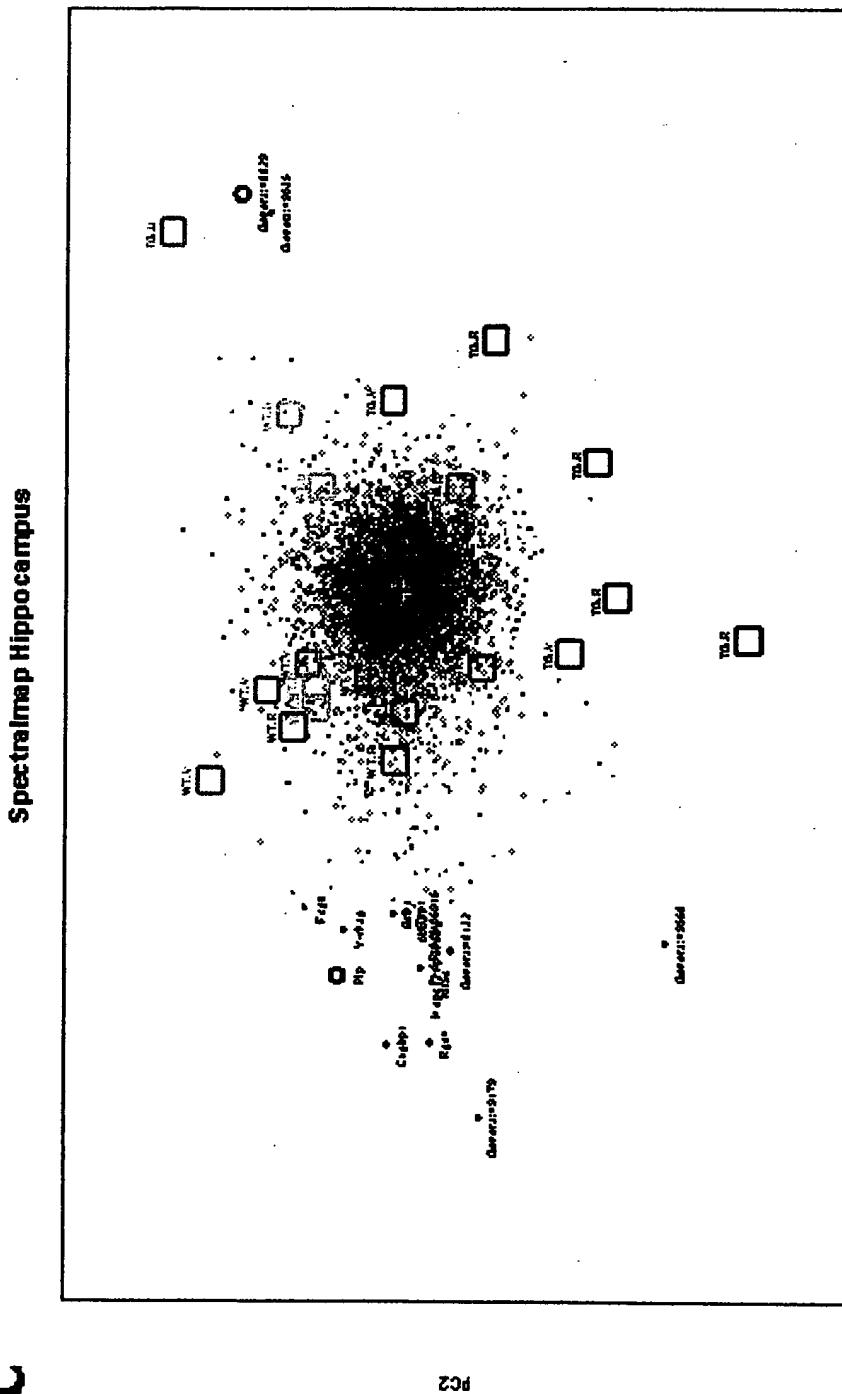


B

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Fig.2 continued



REQUIRES = 25% + 13%, Closure = none, Center = double, Norm = global, Scale = wic, R00 = mean, C01 = mean

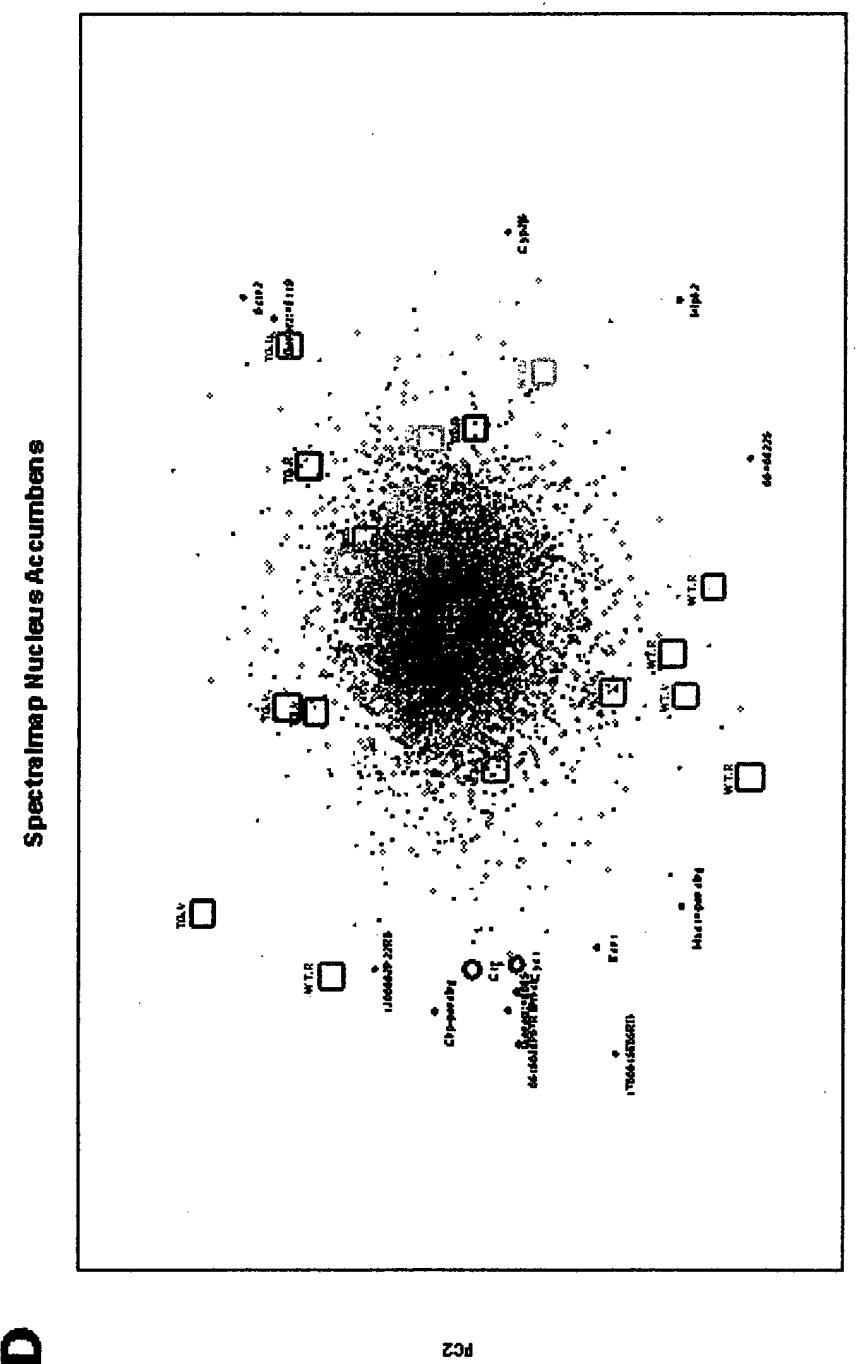
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Fig.2 continued

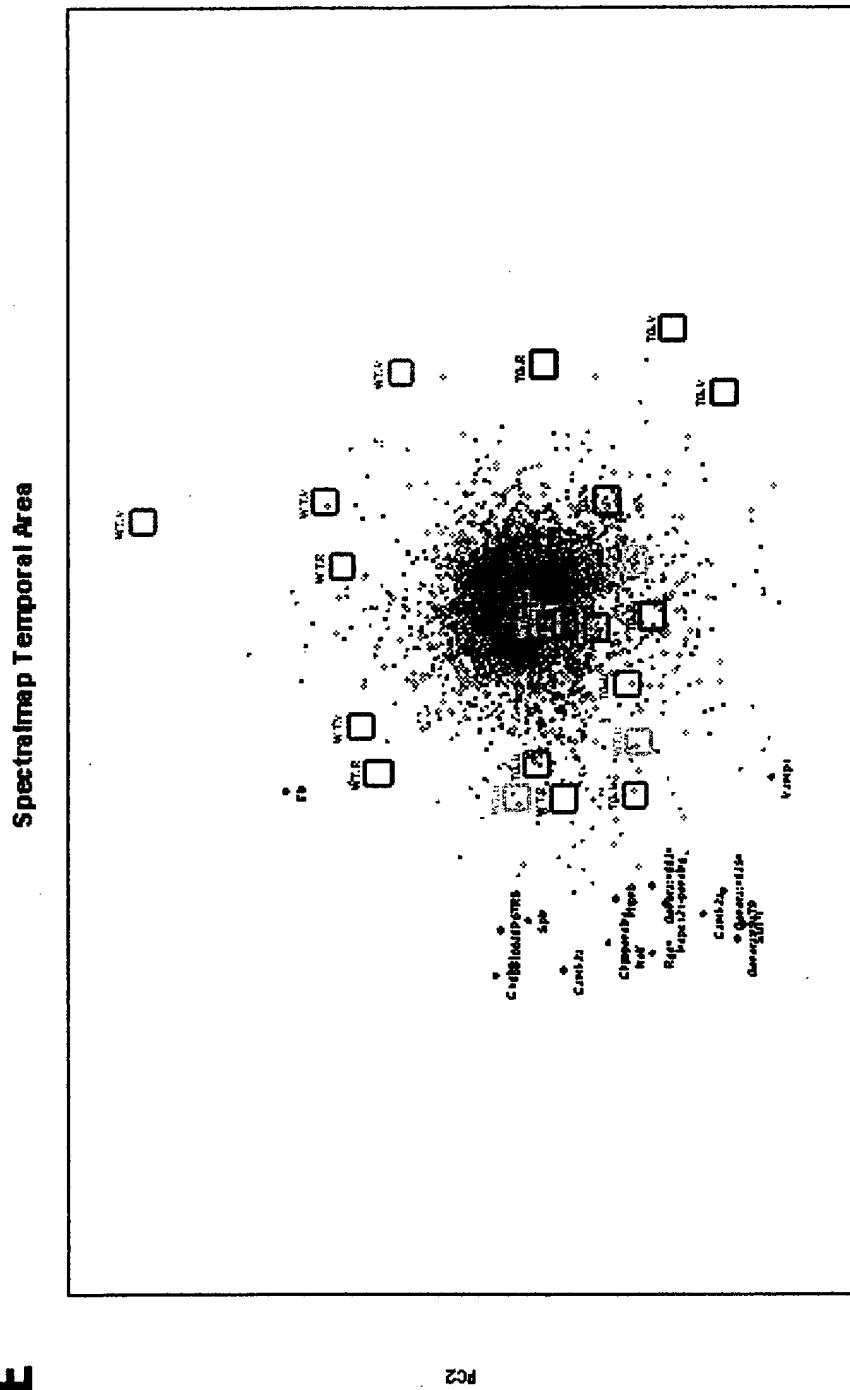


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Fig.2 continued



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Fig.2 continued

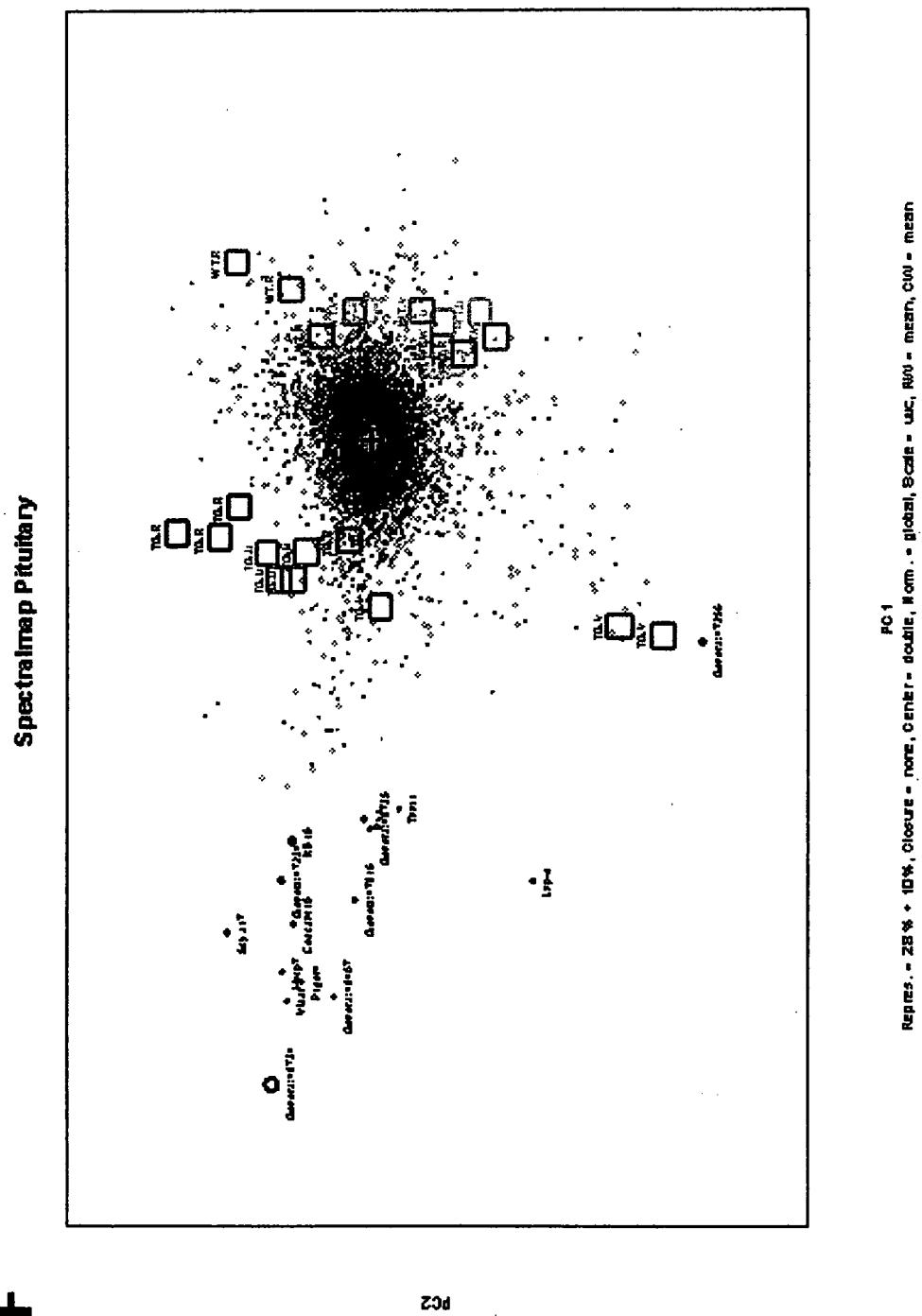
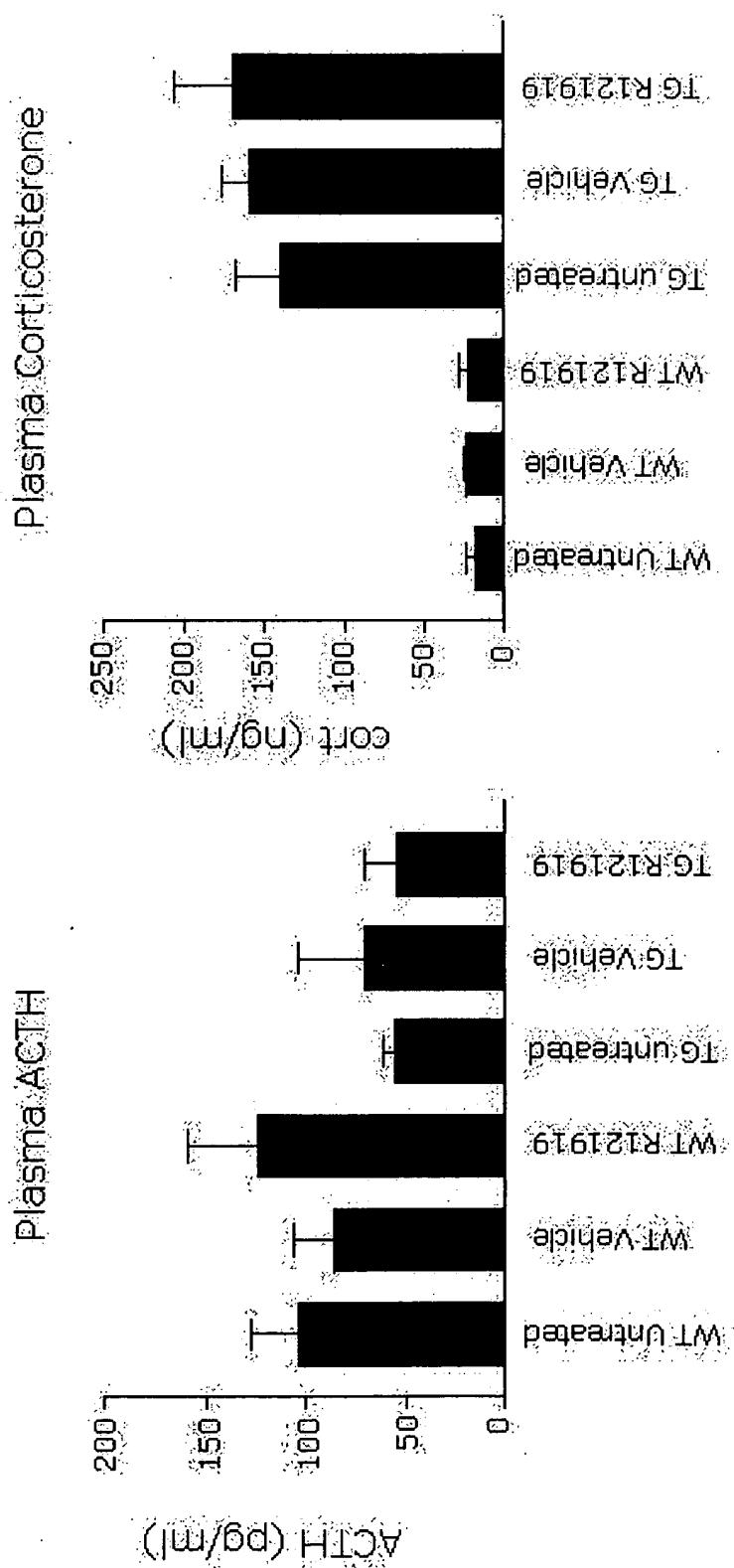


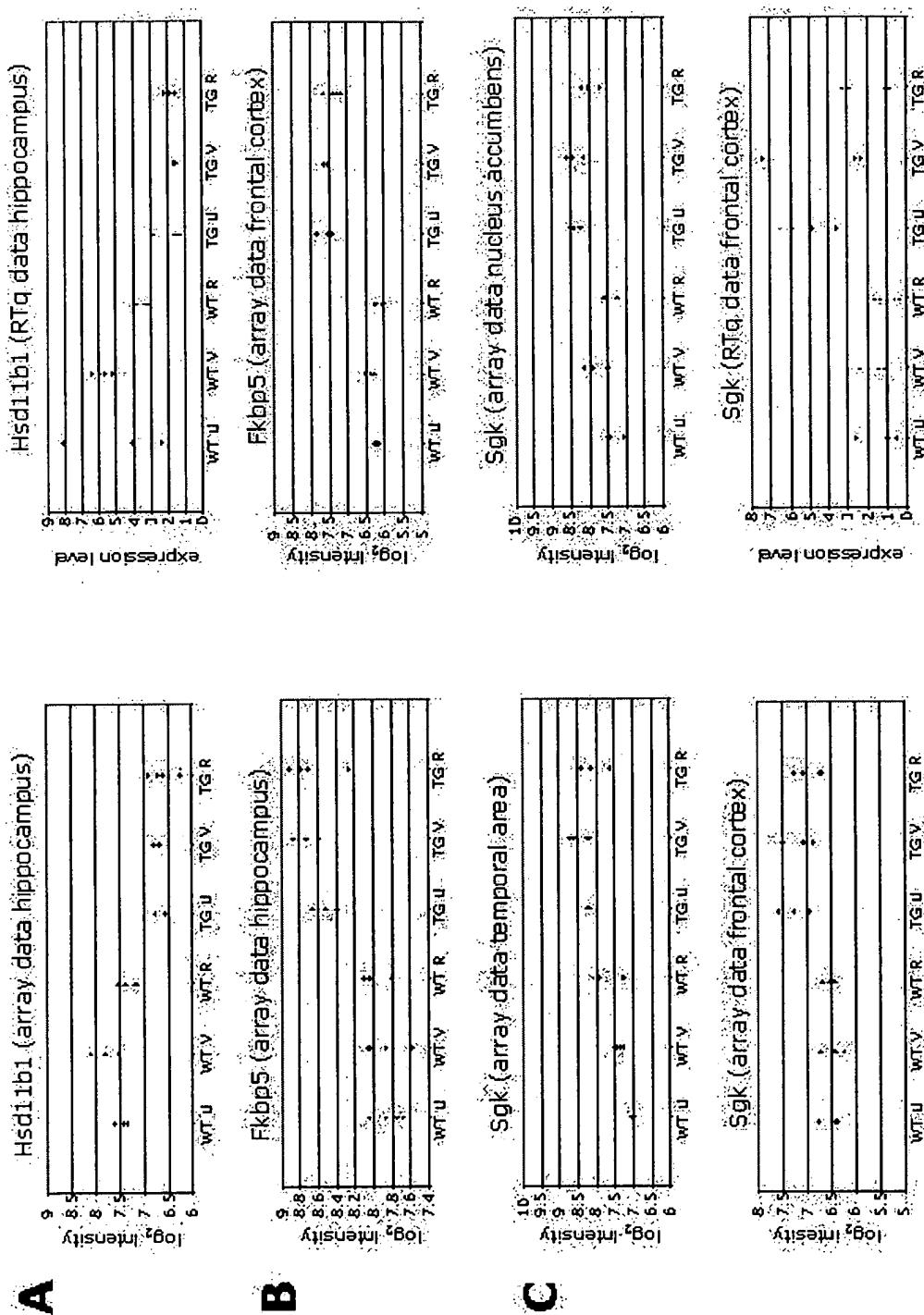
Fig. 3



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Fig. 4



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Fig.5

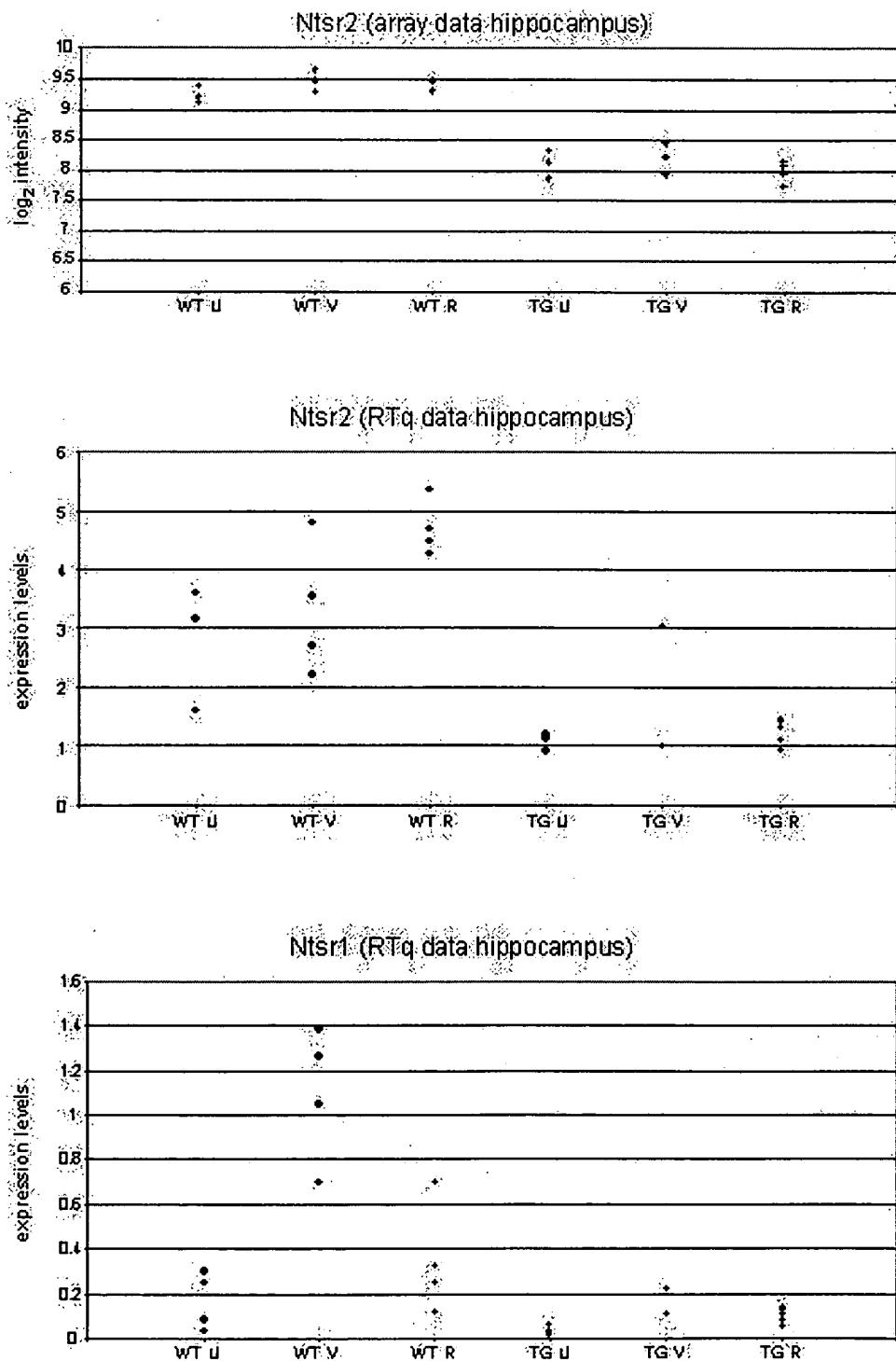


Fig.6

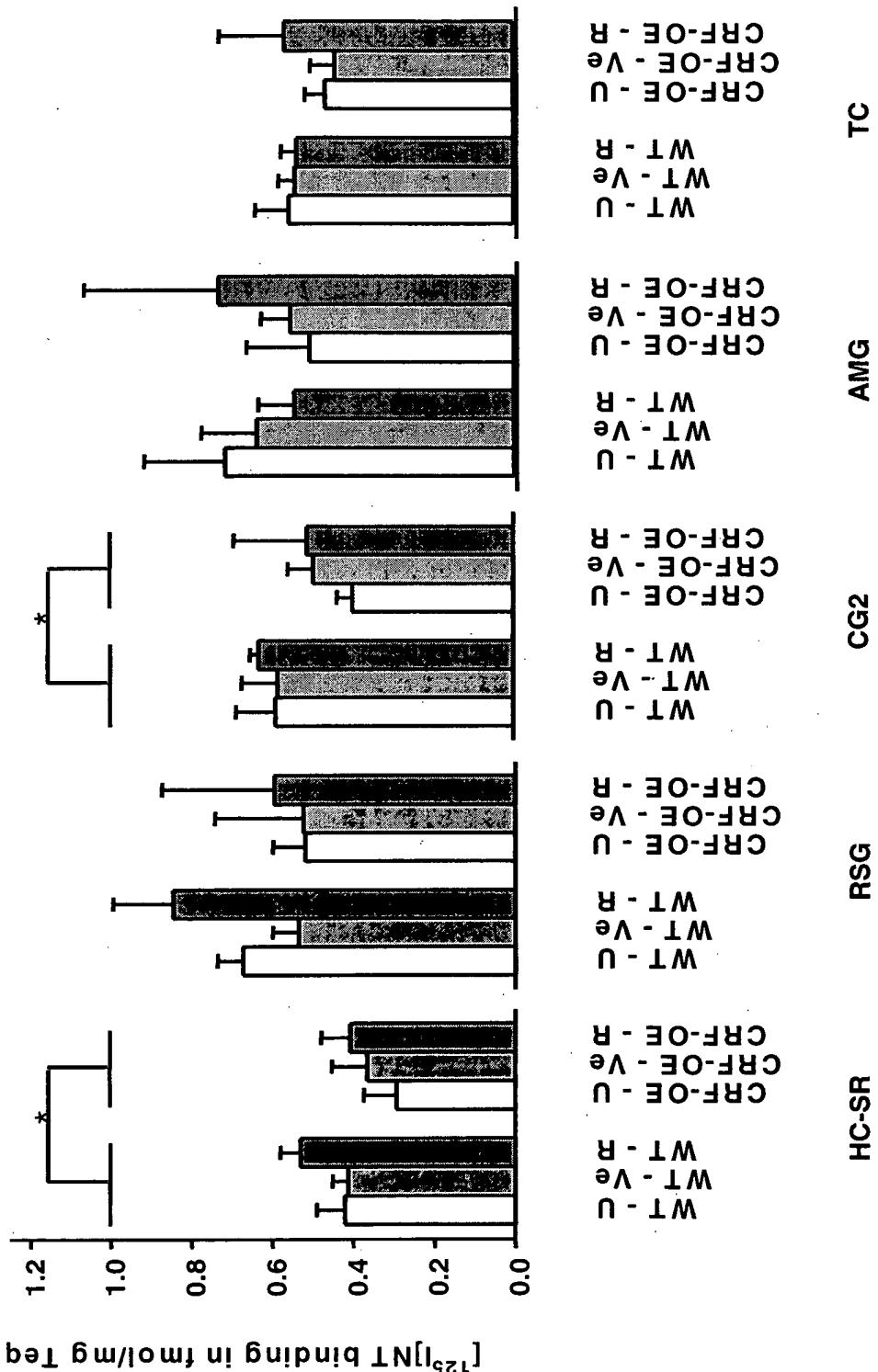
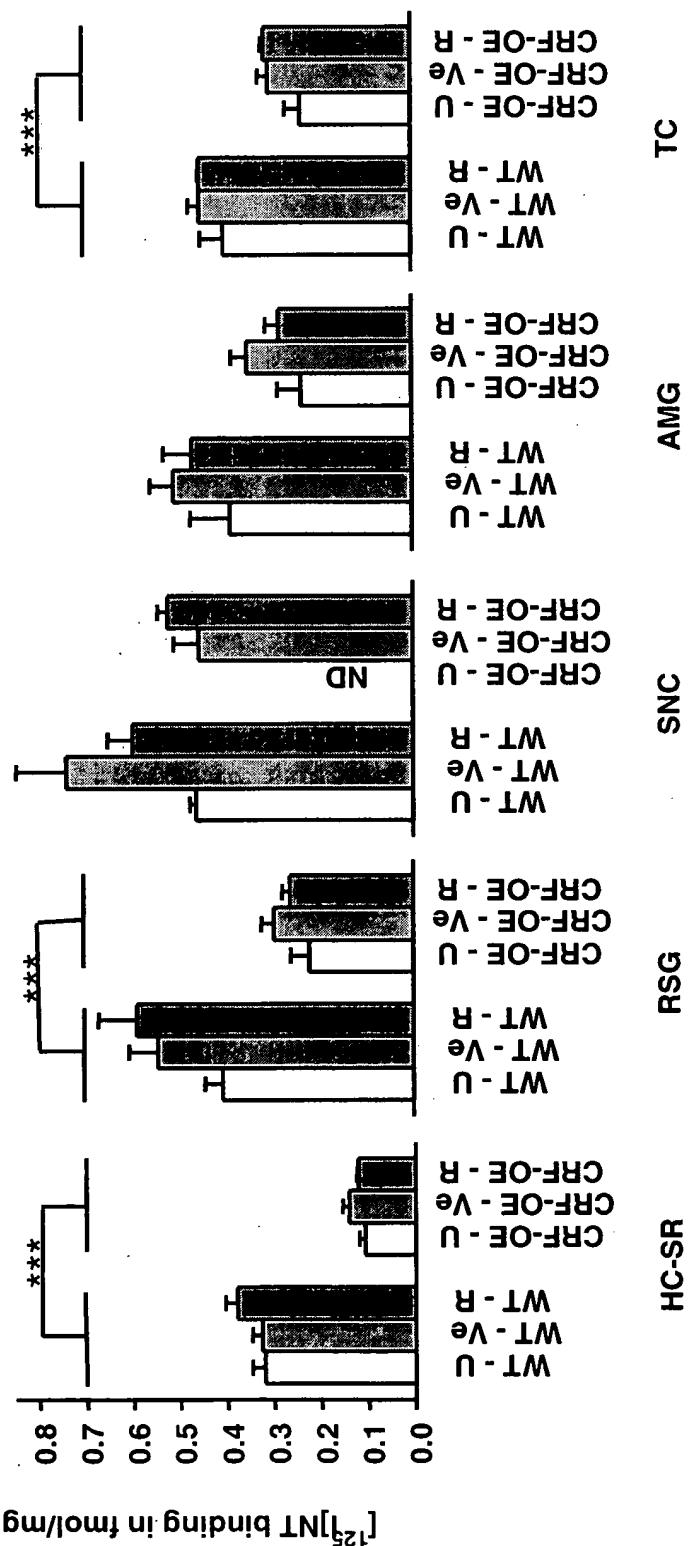


Fig.7



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